
Sequence Listing was accepted.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)

217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: [year=2008; month=3; day=19; hr=13; min=41; sec=43; ms=274;]

Validated By CRFValidator v 1.0.3

Application No: 10705300 Version No: 3.1

Input Set:

Output Set:

Started: 2008-03-19 13:40:30.679

Finished: 2008-03-19 13:40:32.486

Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 807 ms

Total Warnings: 0

Total Errors: 2

No. of SeqIDs Defined: 391

Actual SeqID Count: 391

Error code Error Description

E 104 Command to process tag does not exist: Tag: 2_ii_D

E 104 Command to process tag does not exist: Tag: 2_ii_D

SEQUENCE LISTING

- (1) GENERAL INFORMATION:
 - (i) APPLICANT: PARMA, et al.
 - (ii) TITLE OF INVENTION: HIGH AFFINITY NUCLEIC ACID LIGANDS
 TO LECTINS
 - (iii) NUMBER OF SEQUENCES: 391
 - (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Swanson & Bratschun, L.L.C.
 - (B) STREET: 8400 E. Prentice Avenue, Suite 200
 - (C) CITY: Englewood
 - (D) STATE: Colorado
 - (E) COUNTRY: USA
 - (F) ZIP: 80111
 - (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Diskette, 3 1/2 diskette, 1.44 MB
 - (B) COMPUTER: IBM pc compatible
 - (C) OPERATING SYSTEM: MS-DOS
 - (D) SOFTWARE: WordPerfect 6.0
 - (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: 10705300
 - (B) FILING DATE: 10-NOV-2003
 - (C) CLASSIFICATION: <Unknown>
 - (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: US/08/952,793
 - (B) FILING DATE: 20-NOV-1997
 - (A) APPLICATION NUMBER: PCT/US96/09455
 - (B) FILING DATE: 05-JUN-1995
 - (A) APPLICATION NUMBER: 08/479,724
 - (B) FILING DATE: 07-JUN-1995
 - (A) APPLICATION NUMBER: 08/472,256
 - (B) FILING DATE: 07-JUN-1995
 - (A) APPLICATION NUMBER: 08/472,255
 - (B) FILING DATE: 07-JUN-1995
 - (A) APPLICATION NUMBER: 08/477,829
 - (B) FILING DATE: 07-JUN-1995
 - (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Barry J. Swanson
 - (B) REGISTRATION NUMBER: 33,215
 - (C) REFERENCE/DOCKET NUMBER: NEX40C/PCT
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: (303) 793-3333
 - (B) TELEFAX: (303) 793-3433
- (2) INFORMATION FOR SEQ ID NO: 1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 98 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: RNA
 - (ix) FEATURE:
 - (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
 - (ix) FEATURE:
 - (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

GGGAAAAGCG	AAUCAUACAC	AAGANNNNNN	иииииииии	иииииииии	50
имимимими	ииииииииии	NNNNGCUCCG	CCAGAGACCA	ACCGAGAA	98

	141414	ALOADOA AOOAGAGO COO CAGAGAGA ACCAGAGA	50
(2)	INFO	RMATION FOR SEQ ID NO: 2:	
` ,		SEQUENCE CHARACTERISTICS:	
	` '	(A) LENGTH: 41 base pairs	
		(B) TYPE: nucleic acid	
		(C) STRANDEDNESS: single	
		(D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: RNA	
		FEATURE:	
	(— - · · /	(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine	
	(ix)	FEATURE:	
		(D) OTHER INFORMATION: All U's are 2'-NH2 uracil	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 2:	
		UACGACU CACUAUAGGG AAAAGCGAAU CAUACACAAG A	41
(2)	INFO	RMATION FOR SEQ ID NO: 3:	
	(i)	SEQUENCE CHARACTERISTICS:	
		(A) LENGTH: 24 base pairs	
		(B) TYPE: nucleic acid	
		(C) STRANDEDNESS: single	
		(D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: RNA	
	(ix)	FEATURE:	
		(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine	
	(ix)	FEATURE:	
		(D) OTHER INFORMATION: All U's are 2'-NH2 uracil	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 3:	
	UUC	UCGGUUG GUCUCUGGCG GAGC	24
(2)	TNFO	RMATION FOR SEQ ID NO: 4:	
(- /		SEQUENCE CHARACTERISTICS:	
	(-/	(A) LENGTH: 96 base pairs	
		(B) TYPE: nucleic acid	
		(C) STRANDEDNESS: single	
		(D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: RNA	
	` '	FEATURE:	
	(± ± * /	(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine	
	(ix)	FEATURE:	
	((D) OTHER INFORMATION: All U's are 2'-NH2 uracil	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 4:	
		AAAAGCG AAUCAUACAC AAGAAUGGUU GGCCUGGGCG CAGGCUUCGA	50
		CUCGGCG GGAACGGGAA UGGCUCCGCC AGAGACCAAC CGAGAA	96
	11011		,
(2)	INFO	RMATION FOR SEQ ID NO: 5:	
	(i)	SEQUENCE CHARACTERISTICS:	
		(A) LENGTH: 98 base pairs	
		(B) TYPE: nucleic acid	
		(C) STRANDEDNESS: single	
		(D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: RNA	
		FEATURE:	
		(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine	
		<u> </u>	

(D) OTHER INFORMATION: All U's are 2'-NH2 uracil

(ix) FEATURE:

	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5: GGGAAAAGCG AAUCAUACAC AAGACAGGCA CUGAAAACUC GGCGGGAACG AAAGUAGUGC CGACUCAGAC GCGUGCUCCG CCAGAGACCA ACCGAGAA	50 98
(2)	INFORMATION FOR SEQ ID NO: 6: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 91 base pairs (B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: RNA	
	(ix) FEATURE:	
	(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine	
	(ix) FEATURE:	
	(D) OTHER INFORMATION: All U's are 2'-NH2 uracil	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:	
	GGGAAAAGCG AAUCAUACAC AAGAAGUCUG GCCAAAGACU CGGCGGGAAC	50
	GUAAAACGGC CAGAAUUGCU CCGCCAGAGA CCAACCGAGA A	91
(2)	INFORMATION FOR SEQ ID NO: 7:	
(- /	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 94 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: RNA	
	(ix) FEATURE:	
	(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine (ix) FEATURE:	
	(D) OTHER INFORMATION: All U's are 2'-NH2 uracil	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:	
	GGGAAAAGCG AAUCAUACAC AAGAGUAGGA GGUUCCAUCA CCAGGACUCG	50
	GCGGGAACGG AAGGUGAUGS GCUCCGCCAG AGACCAACCG AGAA	94
(2)	· · · · · · · · · · · · · · · · · · ·	
	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 95 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: RNA	
	(ix) FEATURE:	
	(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine	
	(ix) FEATURE:	
	(D) OTHER INFORMATION: All U's are 2'-NH2 uracil (xi) SEOUENCE DESCRIPTION: SEO ID NO: 8:	
	GGGAAAAGCG AAUCAUACAC AAGAACAAGG AUCGAUGGCG AGCCGGGGAG	50
	GGCUCGGCGG GAACGAAAUC UGCUCCGCCA GAGACCAACC GAGAA	95
(2)	INFORMATION FOR SEQ ID NO: 9:	
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	(A) LENGTH: 97 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: RNA	

(ix) FEATURE:

	(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine	
	(ix) FEATURE:	
	(D) OTHER INFORMATION: All U's are 2'-NH2 uracil	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:	
	GGGAAAAGCG AAUCAUACAC AAGAUUGGGC AGGCAGAGCG AGACCGGGGG	50
	CUCGGCGGGA ACGGAACAGG AAUGCUCCGC CAGAGACCAA CCGAGAA	97
(2)	INFORMATION FOR SEQ ID NO: 10:	
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 97 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: RNA	
	(ix) FEATURE:	
	(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine	
	(ix) FEATURE:	
	(D) OTHER INFORMATION: All U's are 2'-NH2 uracil	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:	
	GGGAAAAGCG AAUCAUACAC AAGAAAGGGA UGGGAUUGGG ACGAGCGGCC	50
	AAGACUCGGC GGGAACGAAG GGUGCUCCGC CAGAGACCAA CCGAGAA	97
(2)	INFORMATION FOR SEQ ID NO: 11:	
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 96 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: RNA	
	(ix) FEATURE:	
	(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine	
	(ix) FEATURE:	
	(D) OTHER INFORMATION: All U's are 2'-NH2 uracil	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:	
	GGGAAAAGCG AAUCAUACAC AAGACUCGGC GGGAACGAAA GUGUCAUGGU	50
	AGCAAGUCCA AUGGUGGACU CUGCUCCGCC AGAGACCAAC CGAGAA	96
(2)	INFORMATION FOR SEQ ID NO: 12:	
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 98 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: RNA	
	(ix) FEATURE:	
	(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine	
	(ix) FEATURE:	
	(D) OTHER INFORMATION: All U's are 2'-NH2 uracil	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:	
	GGGAAAAGCG AAUCAUACAC AAGACUCGGC GGGAACGUGA AGUGGGUAGG	50
	UAGCUGAAGA CGGUCUGGGC GCCAGCUCCG CCAGAGACCA ACCGAGAA	98
(0)	TAMORNA MARKA TON TON TO TO TO TO	
(2)	INFORMATION FOR SEQ ID NO: 13:	
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 99 base pairs	
	(B) TYPE: nucleic acid	

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: RNA	
(ix) FEATURE:	
(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine	
(ix) FEATURE:	
(D) OTHER INFORMATION: All U's are 2'-NH2 uracil	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:	
GGGAAAAGCG AAUCAUACAC AAGAAAGGGA UGGGAUUGGG ACGAGCGGCC	50
AAGACUCGGC GGGAACGAAG GGUCCGCUCC GCCAGAGACC AACCGAGAA	99
INFORMATION FOR SEQ ID NO: 14:	
(i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 98 base pairs	
(B) TYPE: nucleic acid	
(C) STRANDEDNESS: single	
(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: RNA	
(ix) FEATURE:	
(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine	
	50
ACGAUCACUU GGUACUAAAA GCCCGCUCCG CCAGAGACCA ACCGAGAA	98
INFORMATION FOR SEC ID NO. 15.	
` '	
-	
(ii) MOLECULE TYPE: RNA	
(ix) FEATURE:	
(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine	
(ix) FEATURE:	
(D) OTHER INFORMATION: All U's are 2'-NH2 uracil	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15:	
GGGAAAAGCG AAUCAUACAC AAGACUCGGC GGGAAUCGAA AGUGUACUGA	50
AUUAGAACGG UGGGCCUGCU CAUCGUGCUC CGCCAGAGAC CAACCGAGAA	100
INFORMATION FOR SEQ ID NO: 16:	
(i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 103 base pairs	
(B) TYPE: nucleic acid	
(C) STRANDEDNESS: single	
(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: RNA	
(ix) FEATURE:	
•	
	E 0
	50
	100
UMM	103
	(IX) FEATURE: (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine (IX) FEATURE: (D) OTHER INFORMATION: All U's are 2'-NH2 uracil (XI) SEQUENCE DESCRIPTION: SEQ ID NO: 13: GGGAAAAGCG AAUCAUACAC AAGAAAGGGA UGGGAUUGGG ACGAGCGCC AAGACUCGGC GGGAACGAAG GGUCCGCUCC GCCAGAGACC AACCGAGAA INFORMATION FOR SEQ ID NO: 14: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 98 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: RNA (ix) FEATURE: (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine (ix) FEATURE: (D) OTHER INFORMATION: All U's are 2'-NH2 uracil (XI) SEQUENCE DESCRIPTION: SEQ ID NO: 14: GGGAAAAGCG AAUCAUCACA AAGACUCGGC GGGAACGAAG UGUUGAGUA ACGAUCACUU GGUACUAAAA GCCCGCUCCG CCAGAGACCA ACCGAGAA INFORMATION FOR SEQ ID NO: 15: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 100 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: RNA (ix) FEATURE: (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine (ix) FEATURE: (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine (ix) FEATURE: (D) OTHER INFORMATION: All U's are 2'-NH2 uracil (XI) SEQUENCE DESCRIPTION: SEQ ID NO: 15: GGGAAAAGCG AAUCAUACAC AAGACUCGGC GGGAAUCGAA AGUGUACUGA AUUAGAACGG UGGCCUGCU CAUCGUGCUC CGCCAGAGAC CAACCGAGAA INFORMATION FOR SEQ ID NO: 16: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 103 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: RNA

(i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 98 base pairs	
(B) TYPE: nucleic acid	
(C) STRANDEDNESS: single	
(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: RNA	
(ix) FEATURE:	
(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine	
(ix) FEATURE:	
(D) OTHER INFORMATION: All U's are 2'-NH2 uracil	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 17:	
GGGAAAAGCG AAUCAUACAC AAGACAGCGG CGGAGUCAGU GAAAGCGUGG	50
GGGGYGCGGG AGGUCUACCC UGACGCUCCG CCAGAGACCA ACCGAGAA	98
INFORMATION FOR SEQ ID NO: 18:	
(i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 95 base pairs	
(B) TYPE: nucleic acid	
(C) STRANDEDNESS: single	
(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: RNA	
(ix) FEATURE:	
(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine	
(ix) FEATURE:	
(D) OTHER INFORMATION: All U's are 2'-NH2 uracil	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 18:	
GGGAAAAGCG AAUCAUACAC AAGACGGCUG UGUGUGGUAG CGUCAUAGUA	50
GGAGUCGUCA CGAACCAAGG CGCUCCGCCA GAGACCAACC GAGAA	95
INFORMATION FOR SEQ ID NO: 19:	
(i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 98 base pairs	
(B) TYPE: nucleic acid	
(C) STRANDEDNESS: single	
(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: RNA	
(ix) FEATURE:	
(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine	
(ix) FEATURE:	
(D) OTHER INFORMATION: All U's are 2'-NH2 uracil	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19:	
GGGAAAAGCG AAUCAUACAC AAGACGGCUG UGUGGUGUUG GAGCGUCAUA	50
GUAGGAGUCG UCACGAACCA AGGCGCUCCG CCAGAGACCA ACCGAGAA	98
THEODMATION FOR GEO. ID. NO. 20.	
INFORMATION FOR SEQ ID NO: 20: (i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 97 base pairs	
(B) TYPE: nucleic acid	
(C) STRANDEDNESS: single	
(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: RNA	
(ix) FEATURE:	
(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine	
(ix) FEATURE:	
(D) OTHER INFORMATION: All U's are 2'-NH2 uracil	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:	
GGGAAAAGCG AAUCAUACAC AAGACGAUGC GAGGCAAGAA AUGGAGUCGU	50

(2)

(2)

(2)

(2)	INFO	RMATION FOR SEQ ID NO: 21:	
	(i)	SEQUENCE CHARACTERISTICS:	
		(A) LENGTH: 95 base pairs	
		(B) TYPE: nucleic acid	
		(C) STRANDEDNESS: single	
		(D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: RNA	
	(ix)	FEATURE:	
	(= ==)	(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine FEATURE:	
	(IX)		
	(\	(D) OTHER INFORMATION: All U's are 2'-NH2 uracil	
		SEQUENCE DESCRIPTION: SEQ ID NO: 21:	50
		AAAAGCG AAUCAUACAC AAGACGUGCG GAGCAAAUAG GGGAUCAUGG	
	AGU	CGUACGA ACCGUUAUCG CGCUCCGCCA GAGACCAACC GAGAA	95
(2)	INFO	RMATION FOR SEQ ID NO: 22:	
	(i)	SEQUENCE CHARACTERISTICS:	
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		(B) TYPE: nucleic acid	
		(C) STRANDEDNESS: single	
		(D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: RNA	
	(ix)	FEATURE:	
		(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine	
	(ix)	FEATURE:	
		(D) OTHER INFORMATION: All U's are 2'-NH2 uracil	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 22:	
	GGG	AAAAGCG AAUCAUACAC AAGACUGGGG AGCAGGAUAU GAGAUGUGCG	50
	GGG	CAAUGGA GUCGUGACGA ACCGCUCCGC CAGAGACCAA CCGAGAA	97
(2)		RMATION FOR SEQ ID NO: 23:	
	(i)	SEQUENCE CHARACTERISTICS:	
		(A) LENGTH: 95 base pairs	
		(B) TYPE: nucleic acid	
		(C) STRANDEDNESS: single	
		(D) TOPOLOGY: linear	
		MOLECULE TYPE: RNA	
	(ix)	FEATURE:	
		(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine	
	(ix)	FEATURE:	
		(D) OTHER INFORMATION: All U's are 2'-NH2 uracil	
		SEQUENCE DESCRIPTION: SEQ ID NO: 23:	
	GGG	AAAAGCG AAUCAUACAC AAGAGUCCGC CCCCAGGGAU GCAACGGGGU	50
	GGC	UCUAAAA GGCUUGGCUA AGCUCCGCCA GAGACCAACC GAGAA	95
(0)			
(2)		DIG T T OIL TO T OIL OIL	
		RMATION FOR SEQ ID NO: 24:	
		SEQUENCE CHARACTERISTICS:	
		SEQUENCE CHARACTERISTICS: (A) LENGTH: 94 base pairs	
		SEQUENCE CHARACTERISTICS: (A) LENGTH: 94 base pairs (B) TYPE: nucleic acid	
		SEQUENCE CHARACTERISTICS: (A) LENGTH: 94 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 94 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 94 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear MOLECULE TYPE: RNA	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 94 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	

(ix) FEATURE:

	(D) OTHER INFORMATION: All U's are 2'-NH2 uracil (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 24: GGGAAAAGCG AAUCAUACAC AAGAGAGAAU GAGCAUGGCC GGGGCAGGAA GUGGGUGGCA ACGGAGGCCA GCUCCGCCAG AGACCAACCG AGAA	50 94
(2)	INFORMATION FOR SEQ ID NO: 25:	
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	(A) LENGTH: 95 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: RNA	
	(ix) FEATURE:	
	(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine	
	(ix) FEATURE:	
	(D) OTHER INFORMATION: All U's are 2'-NH2 uracil	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 25:	
	GGGAAAAGCG AAUCAUACAC AAGAGAUACA GCGCGGGUCU AAAGACCUUG	50
	CCCCUAGGAU GCAACGGGGU GGCUCCGCCA GAGACCAACC GAGAA	95
(2)	INFORMATION FOR SEQ ID NO: 26:	
(2)	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 97 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: RNA	
	(ix) FEATURE:	
	(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine	
	(ix) FEATURE:	
	(D) OTHER INFORMATION: All U's are 2'-NH2 uracil	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 26:	
	GGGAAAAGCG AAUCAUACAC AAGAUGAAGG GUGGUAAGAG AGAGUCUGAG	50
	CUCGUCCUAG GGAUGCAACG GCAGCUCCGC CAGAGACCAA CCGAGAA	97
(2)	THEODWARDON FOR GEO. ID. No. 27.	
(2)		
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 99 base pairs (B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: RNA	
	(ix) FEATURE:	
	(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine	
	(ix) FEATURE:	
	(D) OTHER INFORMATION: All U's are 2'-NH2 uracil	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 27:	
	GGGAAAAGCG AAUCAUACAC AAGACAAACC UGCAGUCGCG CGGUGAAACC	50
	UAGGGUUGCA ACGGUACAUC GCUGUGCUCC GCCAGAGACC AACCGAGAA	99
(2)	INFORMATION FOR SEQ ID NO: 28:	
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 97 base pairs	
	(B) TYPE: nucleic acid (C) STRANDEDNESS: single	
	(C) DIIGHDDHIDD DIIDD DIIDD DIID	

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: RNA

	(ix) FEATURE:	
	(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine	
	(ix) FEATURE:	
	(D) OTHER INFORMATION: All U's are 2'-NH2 uracil	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 28:	
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	ACGUUCCUAG GGAUGCAACG GACGCUCCGC CAGAGACCAA CCGAGAA	97
(2)	INFORMATION FOR SEQ ID NO: 29:	
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 97 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: RNA	
	(ix) FEATURE:	
	(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine	
	(ix) FEATURE:	
	(D) OTHER INFORMATION: All U's are 2'-NH2 uracil	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 29:	
	GGGAAAAGCG AAUCAUACAC AAGAGUGUAC CAAUGGAGGC AAUGCUGCGG	50
	GAAUGGAGGC CUAGGGAUGC AACGCUCCGC CAGAGACCAA CCGAGAA	97
(2)	INFORMATION FOR SEQ ID NO: 30:	
` '	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 97 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: RNA	
	(ix) FEATURE:	
	(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine	
	(ix) FEATURE:	
	(D) OTHER INFORMATION: All U's are 2'-NH2 uracil	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 30:	
	GGGAAAAGCG AAUCAUACAC AAGAGUCCCU AGGGAUGCAA CGGGCAGCAU	50
	UCGCAUAGG	